



#10

SUBSTITUTE SEQUENCE LISTING

<110> PURDUE RESEARCH FOUNDATION

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<120> PSEUDOTYPED RETROVIRUSES AND STABLE CELL LINES FOR THEIR PRODUCTION

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<140> 09/762,224

<141> 2001-02-02

<150> US 60/095,242

<151> 1998-08-04

<150> US 60/112,405

<151> 1998-12-15

<150> PCT/US99/17702

<151> 1999-08-04

<160> 8

<170> PatentIn version 3.0

<210> 1

<211> 2958

<212> DNA

<213> Ross River virus

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<211> 985

<212> PRT

<213> Ross River virus

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 Cys Cys Gly Thr Ser Glu Cys Ser Ser Lys Glu Gln Pro Asp Tyr Gln
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<211> 2224

<212> DNA

<213> Ebola virus

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<211> 676

<212> PRT

<213> Ebola virus

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Gly Pro Cys Ala Gly Asp Phe Ala Phe His Lys Glu Gly Ala Phe Phe
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Leu Tyr Asp Arg Leu Ala Ser Thr Val Ile Tyr Arg Gly Thr Thr Phe
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<211> 2103

<212> DNA

<213> Marburg virus

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<211> 681

<212> PRT

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20      25      30
Val Asp Ser Val Cys Ser Gly Thr Leu Gln Lys Thr Glu Asp Val His
35      40      45

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Leu Met Gly Phe Thr Leu Ser Gly Gln Lys Val Ala Asp Ser Pro Leu
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 Glu Ala Ser Lys Arg Trp Ala Phe Arg Thr Gly Val Pro Pro Lys Asn
 65 70 75 80
 Val Glu Tyr Thr Glu Gly Glu Glu Ala Lys Thr Cys Tyr Asn Ile Ser
 85 90 95
 Val Thr Asp Pro Ser Gly Lys Ser Leu Leu Leu Asp Pro Pro Thr Asn
 100 105 110
 Ile Arg Asp Tyr Pro Lys Cys Lys Thr Ile His His Ile Gln Gly Gln
 115 120 125
 Asn Pro His Ala Gln Gly Ile Ala Leu His Leu Trp Gly Ala Phe Phe
 130 135 140
 Leu Tyr Asp Arg Ile Ala Ser Thr Thr Met Tyr Arg Gly Lys Val Phe
 145 150 155 160
 Thr Glu Gly Asn Ile Ala Ala Met Ile Val Asn Lys Thr Val His Lys
 165 170 175
 Met Ile Phe Ser Arg Gln Gly Gln Gly Tyr Arg His Met Asn Leu Thr
 180 185 190
 Ser Thr Asn Lys Tyr Trp Thr Ser Ser Asn Gly Thr Gln Thr Asn Asp
 195 200 205
 Thr Gly Cys Phe Gly Ala Leu Gln Glu Tyr Asn Ser Thr Lys Asn Gln
 210 215 220
 Thr Cys Ala Pro Ser Lys Ile Pro Pro Pro Leu Pro Thr Ala Arg Pro
 225 230 235 240
 Glu Ile Lys Leu Thr Ser Thr Pro Thr Asp Ala Thr Lys Leu Asn Thr
 245 250 255
 Thr Asp Pro Ser Ser Asp Asp Glu Asp Leu Ala Thr Ser Gly Ser Gly
 260 265 270
 Ser Gly Glu Arg Glu Pro His Thr Thr Ser Asp Ala Val Thr Lys Gln
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 Gly Leu Ser Ser Thr Met Pro Pro Thr Pro Ser Pro Gln Pro Ser Thr
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 Pro Gln Gln Gly Gly Asn Asn Thr Asn His Ser Gln Asp Ala Val Thr
 305 310 315 320
 Glu Leu Asp Lys Asn Asn Thr Thr Ala Gln Pro Ser Met Pro Pro His
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 Asn Thr Thr Thr Ile Ser Thr Asn Asn Thr Ser Lys His Asn Phe Ser
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 Thr Leu Ser Ala Pro Leu Gln Asn Thr Thr Asn Asp Asn Thr Gln Ser
 355 360 365
 Thr Ile Thr Glu Asn Glu Gln Thr Ser Ala Pro Ser Ile Thr Thr Leu
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 Pro Pro Thr Gly Asn Pro Thr Thr Ala Lys Ser Thr Ser Ser Lys Lys

385		390		395		400									
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Asp	Gln	His	Ala	Ser	Pro	Asn	Ile	Ser	Leu	Thr	Leu	Ser	Tyr	Phe	Pro
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28